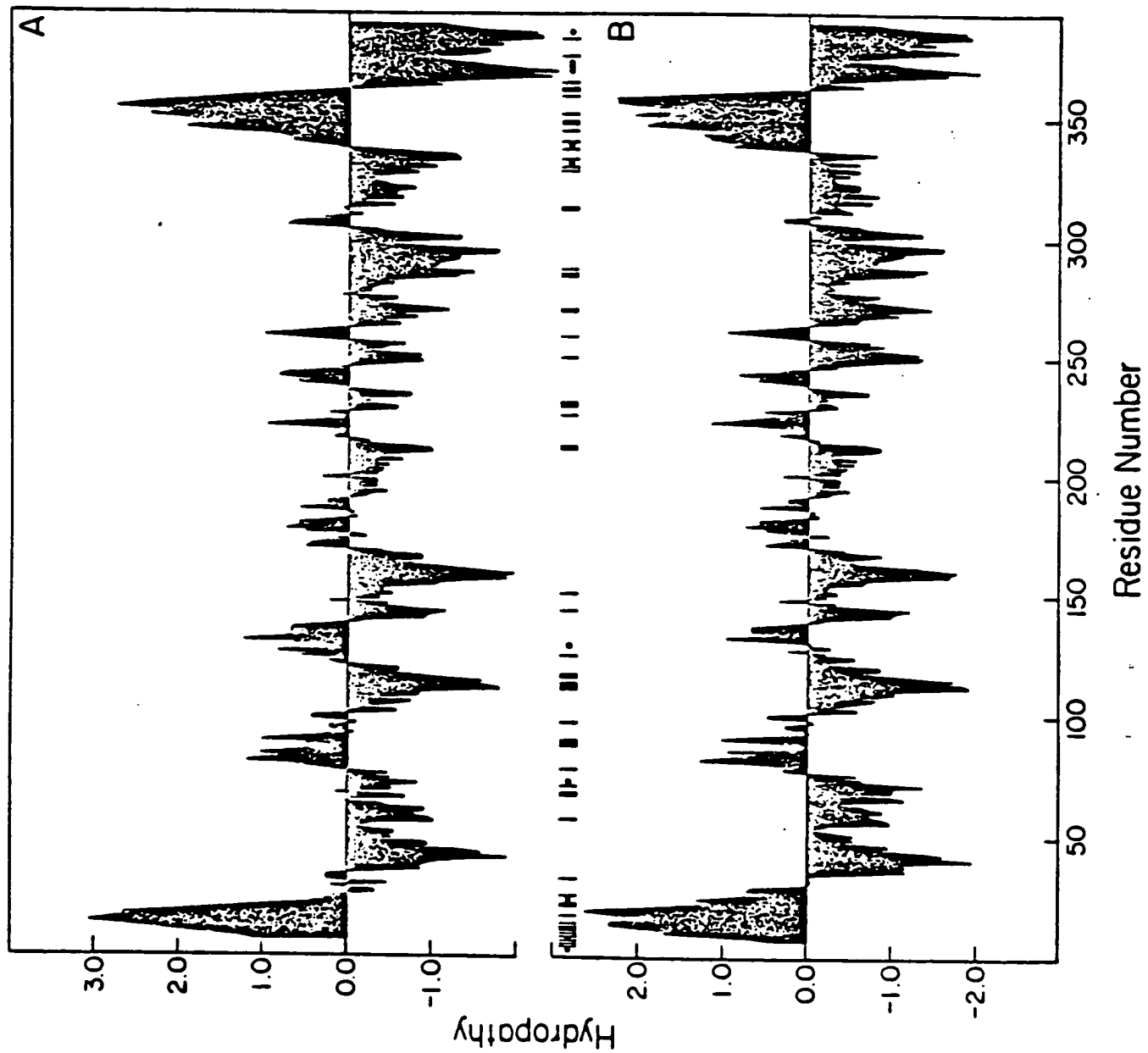


751 GAAACACAGGCGACCGTGGCCCTATACAGCTTAAAAATGCGGGGTGGACAGGCCCCAAGCCCCCGTACACCCAGC
GAGAACCGGCGACCGTGGCCCTATACAGCTTGAAGATCGCGGGGTGGACAGGCCCCAAGCCCCCGTACACCCAGC
751 GluAsnGlnAroThrValAlaValIvrSerLeuLysIleAlaGlyTrpHisGlyProLysProProIvrThrSer
GluAsnGlnAroThrValAlaValIvrSerLeuLysIleAlaGlyTrpHisGlyProLysProProIvrThrSer

FIGURE 1A

FIGURE 2



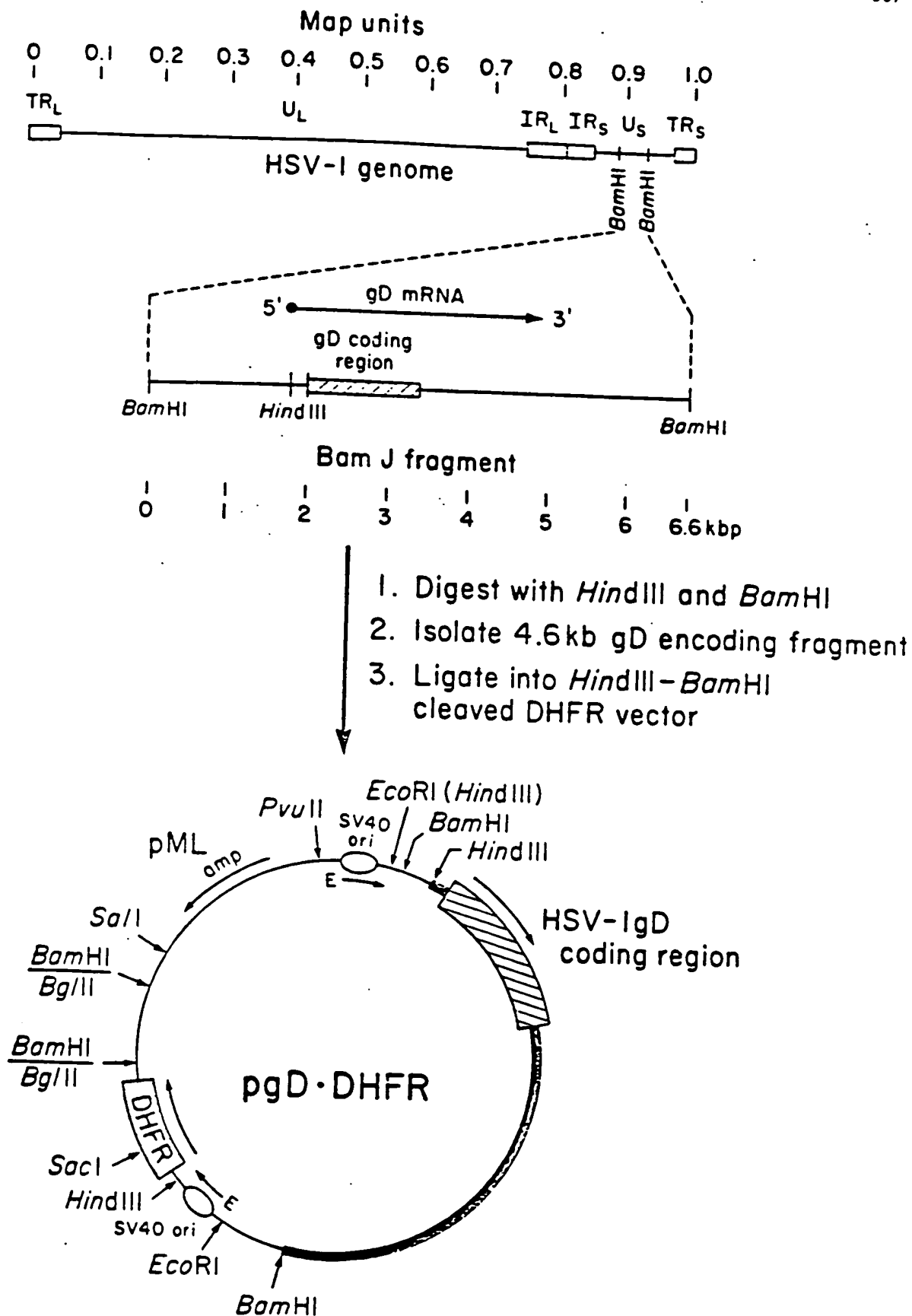


FIGURE 3

A

08/459141



B

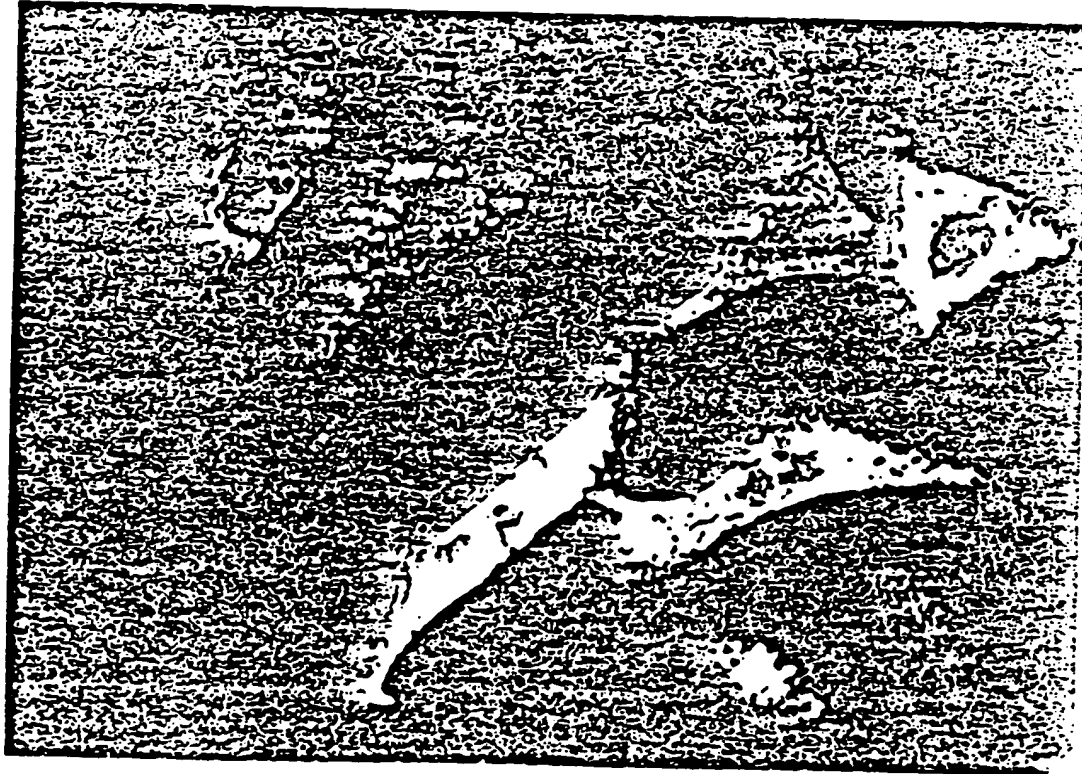


FIGURE 4

A

B

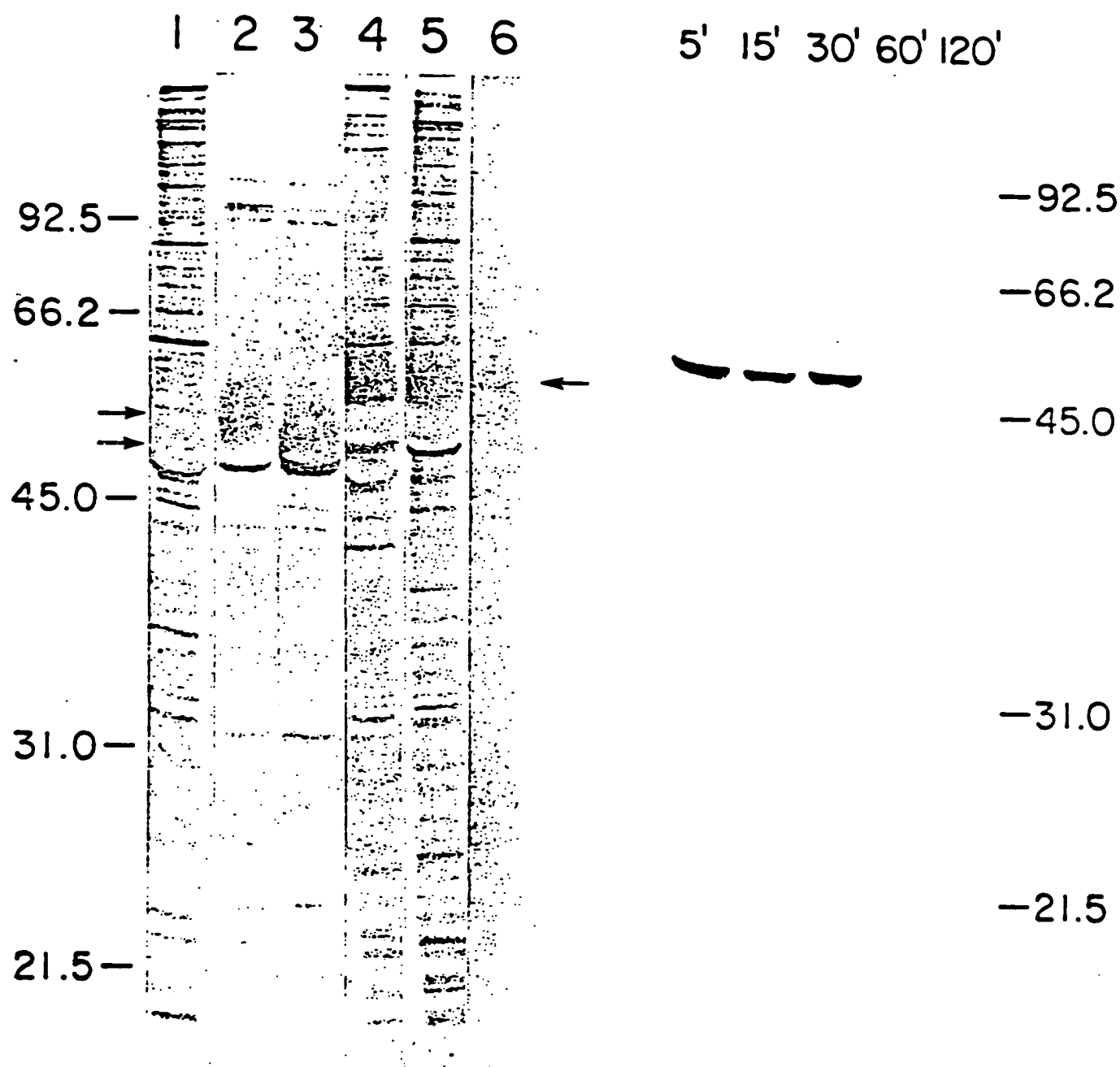


FIGURE 5

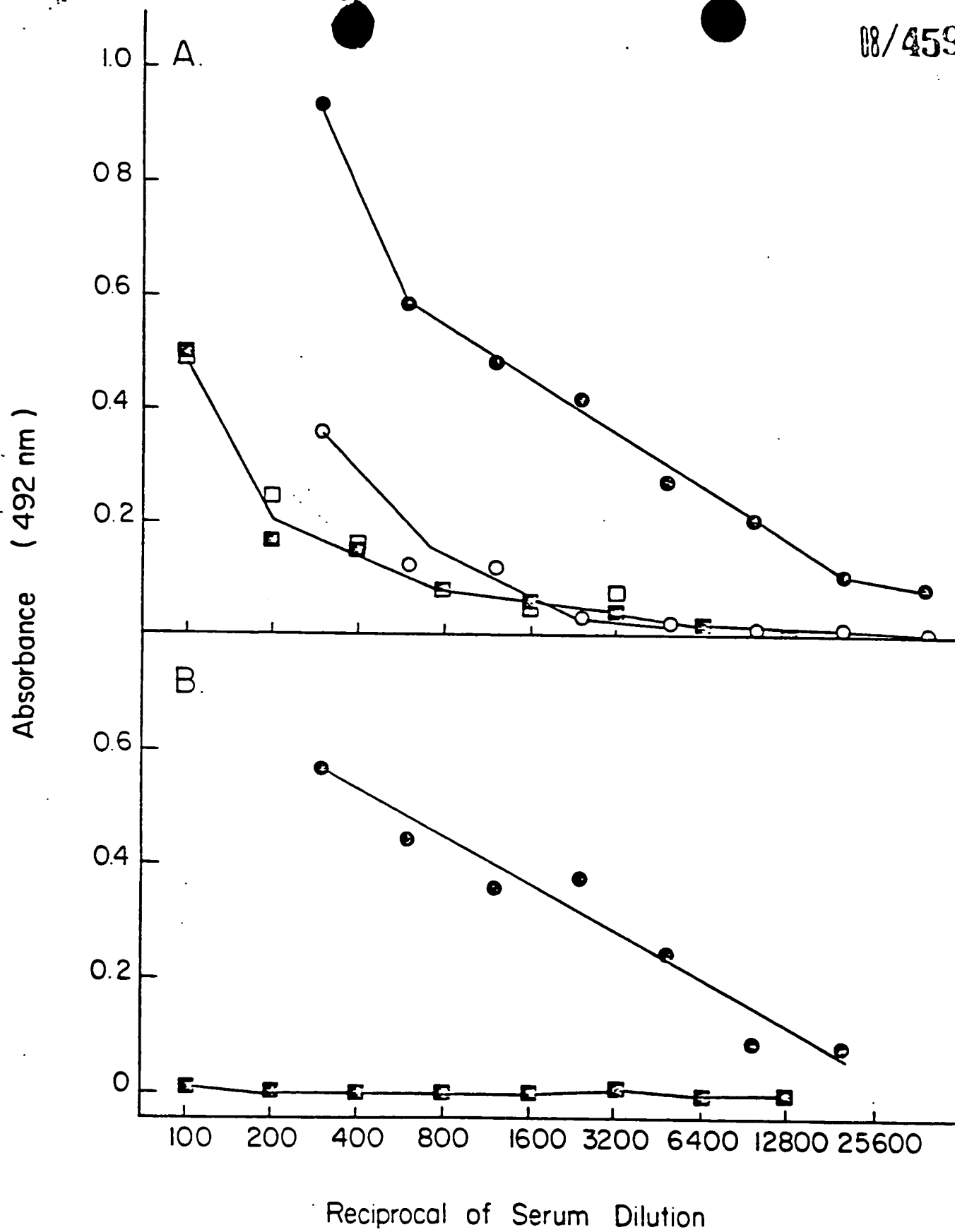


FIGURE 6

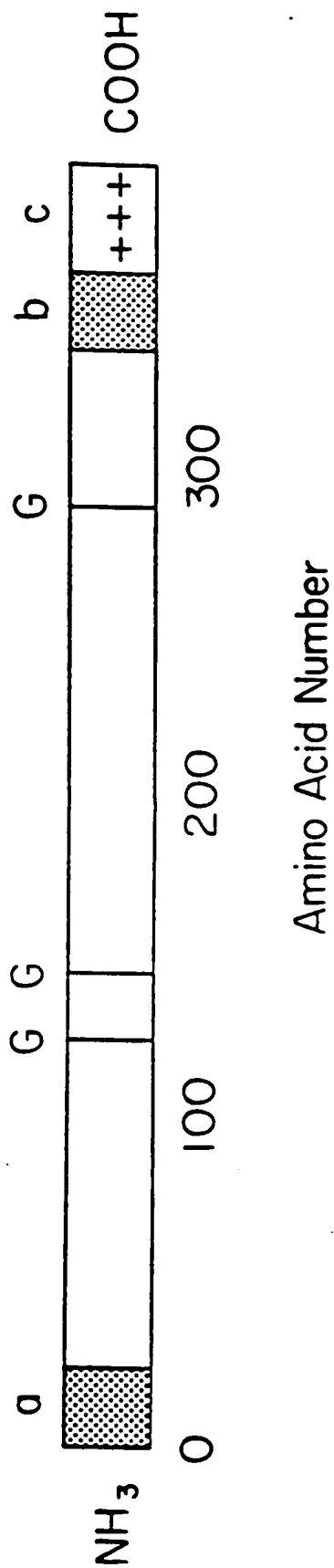


FIGURE 7

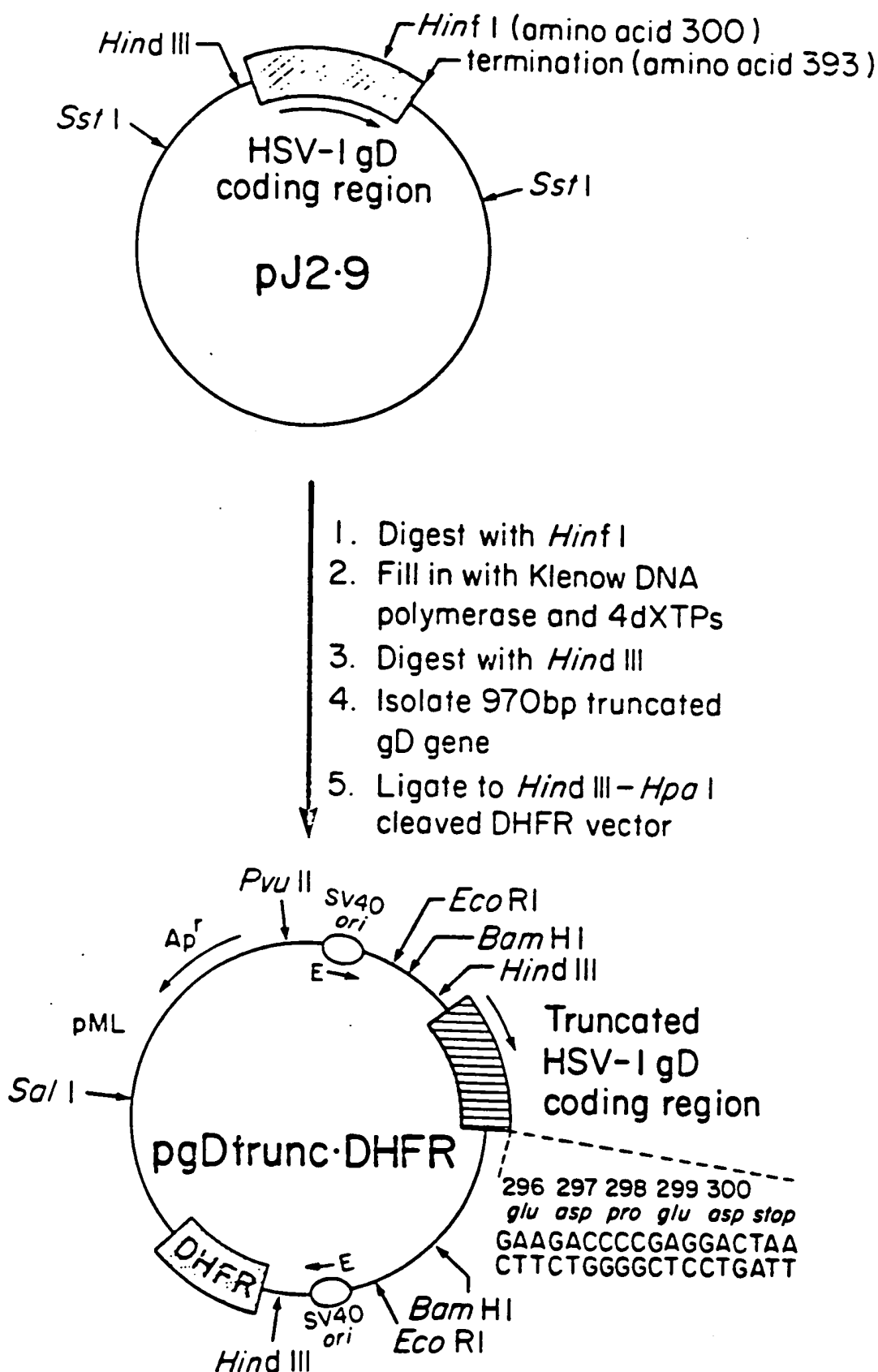


FIGURE 8

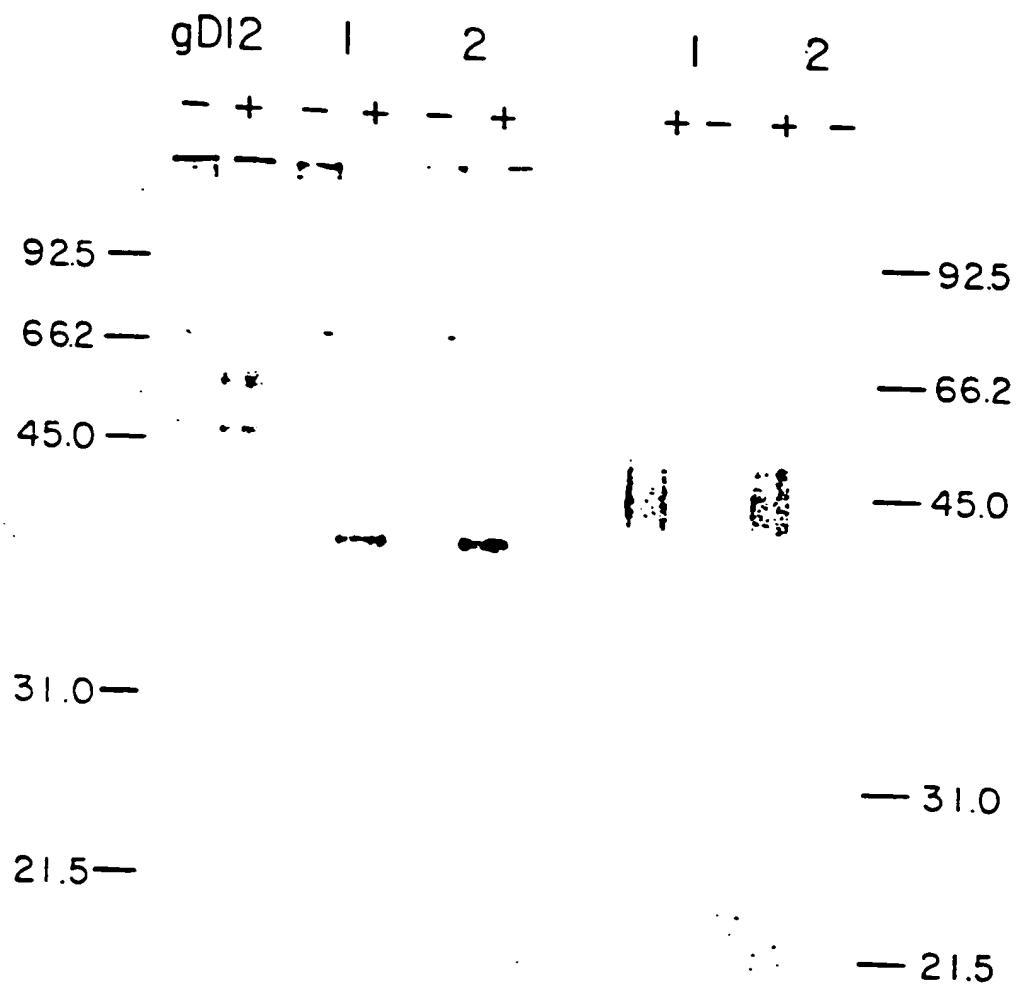


FIGURE 9

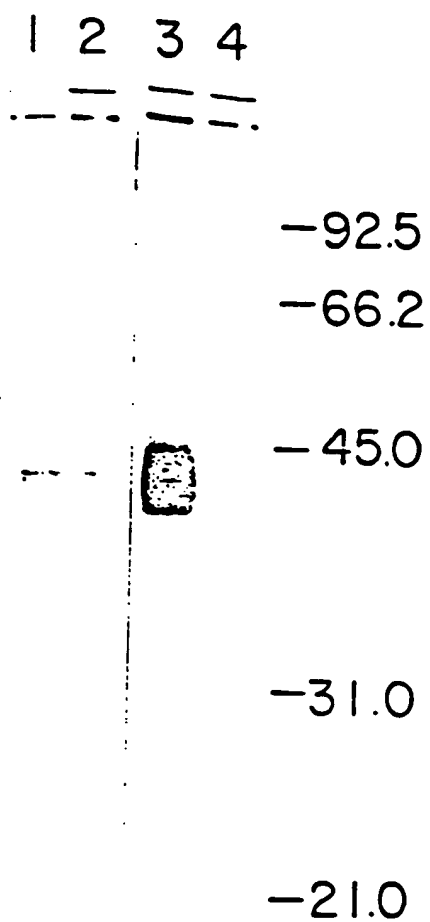


FIGURE 10

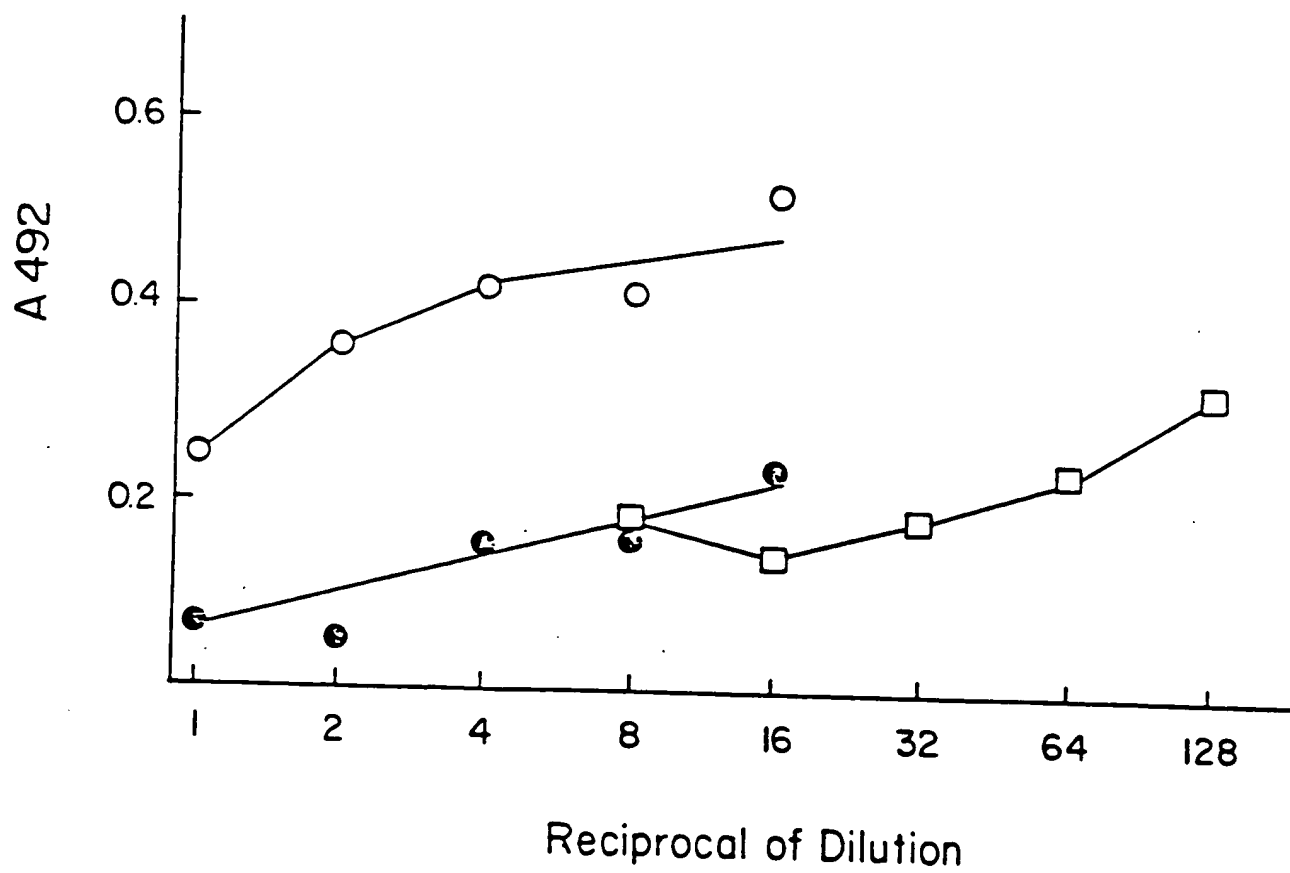


FIGURE 11

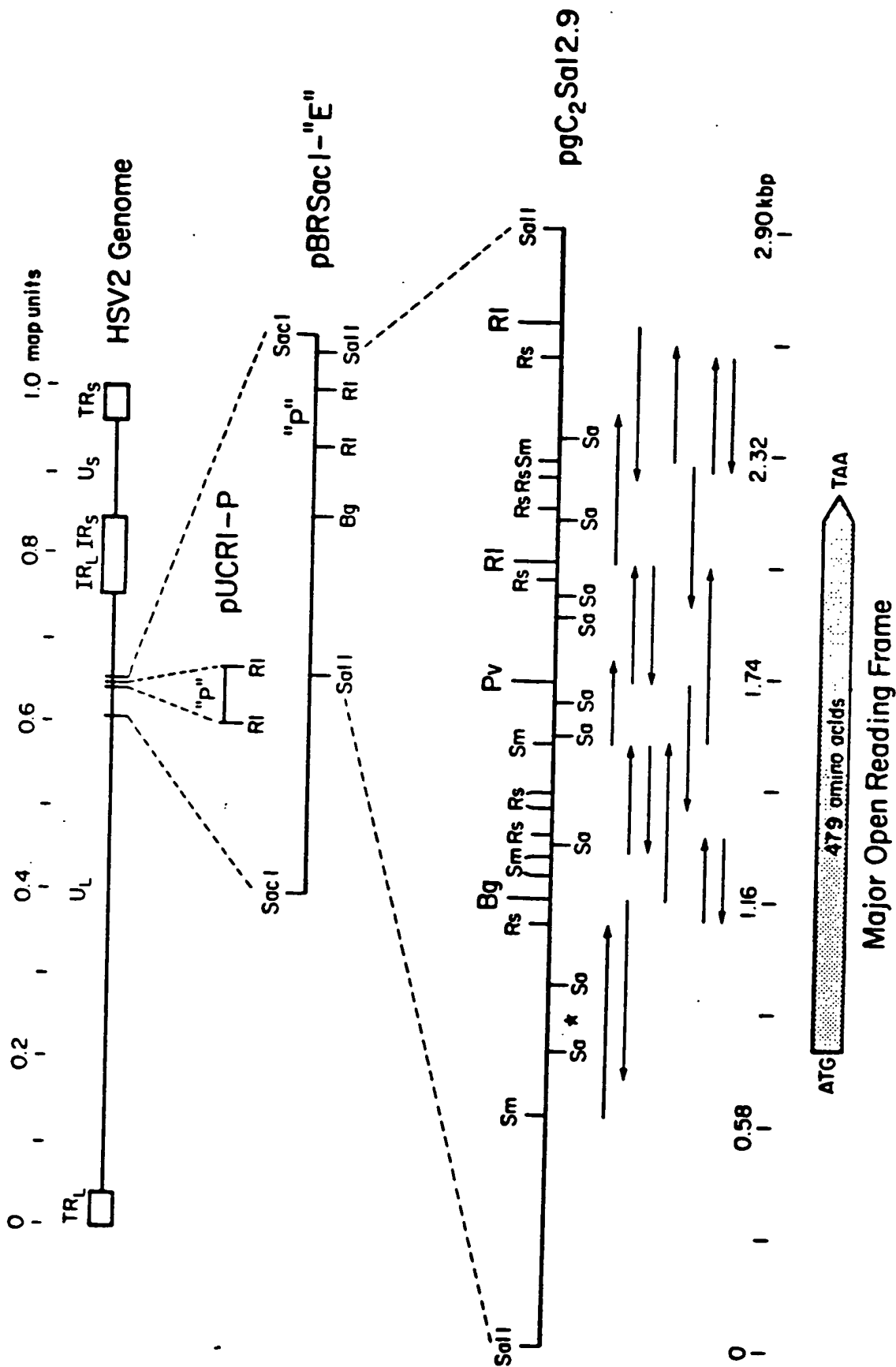


FIGURE 12

[illegible]

MSV-1	GACTGACGTTCTCGGACGCAATGCCACC66GCT66CCCTG66T6CT6LC66G6CA	1320
MSV-2	GACTCCTGTCGTTCTCTCGGCGCAACGCCAGCGGCACGGCATCGGTGCTGCGCGCGGCA	
MSV-1	ACCATCACCATGGAATTTGGGTCGGCATGTGCTGTCACGGCCGGCTGCTGCTCCCGAG	1380
MSV-2	ACCATTACCATGGAGTTTACGGGCGACCATGCGGCTGTCACGGCCGGCTGCTGCTCCCGAG	
MSV-1	GGCGTGACGTTTGCCTGGTTCCTGGGGGACGACCCCTCACC66CGGCTAAGTCGGCCGT	1440
MSV-2	GGGGTGACGTTTGCCTGGTTCCTGGGGGACGACTCTCGCCGGCGGAGAAGGTGGCCGTC	
MSV-1	ACGGCCAGGAGTCTGTCGACACCCCGGGCTGGCTACGGTCCGGTCCACCCTGCCCAT	1500
MSV-2	GGTCCCAGACATCTGTCG66GCGCCCGGCGACCGCCACGATCGGCTCCACCCTGCCGCTC	
MSV-1	TCGTACGACTACAGCGAGTACATCTGCTGGTTGACCGGATATCCGGCCGGGATTCCCGTT	1560
MSV-2	TCGTACGAGCAGACCGAGTACATCTGCGCGCTGGCGGGATACCCGGACGGAATTCGGGTC	
MSV-1	CTAGAGCACCACGGCAGTCAACGACCCCCACCCAGGGAGCCCCACCGAGCGGCAAGTGTATC	1620
MSV-2	CTAGAGCACCACGGCAGCACCACGCCCCCGCGCGGGACCCCCACCGAGCGGCAAGTGTATC	
MSV-1	GAGGCGATCGAGTGGGTGGGGATTGGGAATCGGGGTTCTCGCGGCGGGGGTCTGCTGCTA	1680
MSV-2	CGGGCGGTGGAGGGGGCGGGGATCGGAGTGGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	
MSV-1	ACGGCAATCGTGACGTCGTCGACATCACAGTCGGCGGAGCGTCATCGGGGGTAACGC	1740
MSV-2	ACCGGGTAGTGTACCTCACCCACGCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG	
	MSV-1 qC, MSV-2 qF termination codons	
MSV-1	GAGACCCCCCGTTACCTTTTAAATATCTATAGTTTGGTCCCCCTT---CTATCCCG	1800
MSV-2	CGGGGCGGGCGCCCGGCGCGGCT-TGCTTCTTT-TCCACCCCTTCCGTCGCCCGTACCC	
MSV-1	CC-----CACCCTGGGCGCTATAAAGCC-GCCACCCCTCTC	1860
MSV-2	ACCACACCCACCCACCCCGCGCTGCGCGGGCGTTATAAGC--CGCCGCACTGCG	
	"TATA 2"	
MSV-1	TTCCCTCAGGTC---ATCCTTGGTC-GATCCCGAAGCAGACAGCGGTGGAG---CAAAA	1920
MSV-2	TTTTCCACCGGAAATCCTCGGCGCGATCC-GAACGGCGCAGCGCGGTGGGCTCCAAA	
MSV-1	CGCCTCCCCCTGAGCC-GCTTTCCTACCAACACACCGGCATGCC---T-CT--G-----	1980
MSV-2	CGCCTCGGGAAGAGAGCGCCCGCCCGGAT-ATTCAAGCCCGGGTGGTGTATGGCTTT	
	MSV-2 second open reading frame initiation codon	
MSV-1	-CGGGCATCGGAACAGCC-TACCGGCCCCCTGGGCCCCGGGACACCCCCATGCGGGCTCG	2040
MSV-2	CCGTGCTTCGGGACCCGCTACCAAGCCCCCTGCGCCCGCGGCTCCCCCGGCGGGCTCG	
	730 bp MSV-1 mRNA initiation codon	
MSV-1	GCTCCCGCGCGGCTGGGTTGGCGTCGGGACCATCATCGGGGAGTTGTGATCATTGC	2100
MSV-2	TGTTCCGGCGGTGGCTGGATCGGCGTCGGAGCGATCGTCGGGCGCTTTCGCTGCTGCG	
MSV-1	CGCGTTGGTCTGTCGCTCGCGGCTCGTGGGCACTTTCCTCATGCGACAGCGGATG	2160
MSV-2	CGCGTTGGTCTGTCGACCCCTCGGTCCTCGTGGGACTCTGCGCGTGGACAGCGGCTG	
MSV-1	GCACGAGTTCAACCTCGGGTGCATATCTGGGATCCGACCCCATGGAGCAGGACAGGC	2220
MSV-2	GCAGGAATTCAACCGGGATGCGTGGGTCGGGACCCACCCCGTGGAGCAGGACAGGC	
MSV-1	GGTCGGCGGCTGTAGCGCCCGCGGACCTGATCCCCCGCGGCTGCCAAACAGCTGGC	2280
MSV-2	GGTCGGCGGCTGACGCGCGCGGCGGACCTTATCCCCCGTGGGCGGCGCAAGCAGCTGGC	
MSV-1	CGCCGTCGACGCTCCAGTCGGCAAGATCTCGGGCTACTGGTGGGTGAGCGGAGACGG	2340
MSV-2	CGCTCTGACACGCGTCCAGGCGGAGAGATCTCGGGTACTGGTGGGTGAAACGGAGACGG	
MSV-1	CATTGGGCGCGCTGCGGCTCGTCGACGCGGTTGGCGGATTTGACCACTTTTGGAGGA	2400
MSV-2	CATCCGGACCTGTCTGAGACTCGTCGACAGGCTCAGTGGCATCGACGAGTTTGGAGGA	
MSV-1	GCCCGCCCTTCGATATGCTACTATCCCCGAGTCCCGGGGCTTTGTTGAGTTTGTAAAC	2460
MSV-2	GCTC	
MSV-1	TTCGACCCGCAACGCGCTGGGGCTGCCGTGA	2491

FIGURE 13 (Part 2)

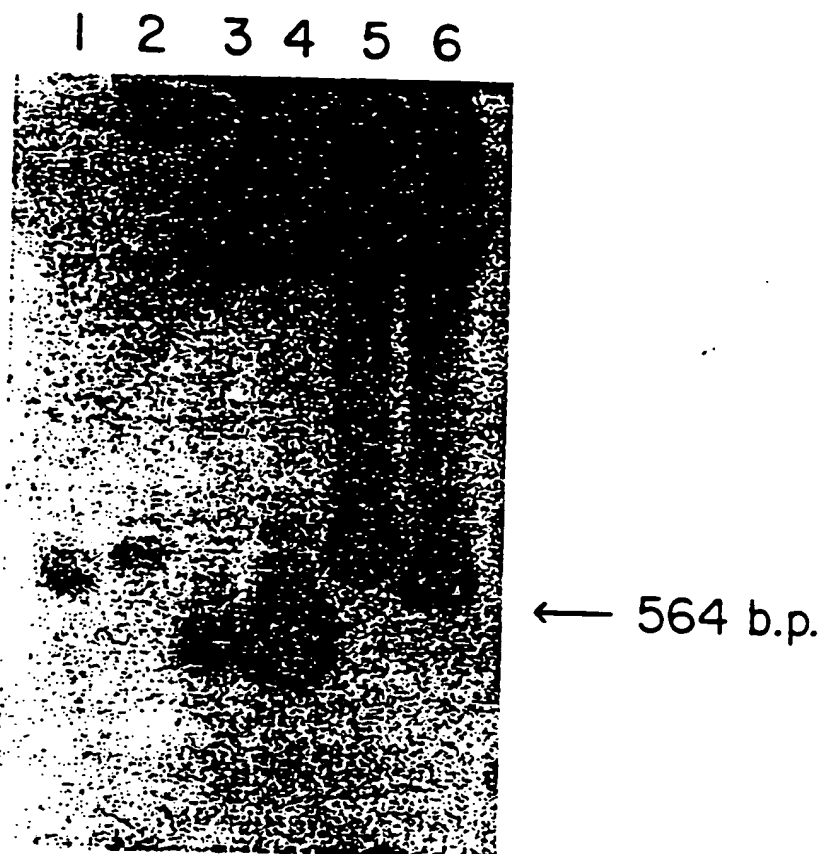


FIGURE 14

A.

```

      C   C   C CC CCN   C N   C CNC   C CC
MSV-1 gC 1  MAPGRVGLAVLVGLLWLGAGVAGGSETASTGPTITAGAVTHAETAPTS
MSV-2 gF 1  MALGRVGLAVLVGLLWVGVVVLA--ASGRTITVGPRA--HA--HAAPS

      NCNC N   N CNC CCNC
MSV-1 gC 50 GSPGSAASPEVTPSTPNP--NVT--NKTTPTEPASPTTPKPTSTPKSPPT
MSV-2 gF 48 -----VPR--NAPRTTTPPPQPRKATKS

      NCNC MN   C   C CCC   N N C
      ....
MSV-1 gC 101 STDPKPK--N--PAKSGRPTKPPG-PV--D--RRDPLARYGSRVQIN--RFR
MSV-2 gF 71 KASTAKAPP--P-KTG--PKTSSEPV--N--RRDPLARYGSRVQIN--RFR

      C C   CCCC CC N   CC CC   N   N
      ....
MSV-1 gC 150 STMEFRLQIWRYSMGPSPIAPAPDLEEVLT--N--APPGLLYVDSAPN
MSV-2 gF 117 STTESRLQIWRATATDAEIGTAPSLEEVN--N--APPGLLYVDSAPN

      C   C N   C   CN   C C CC   NC
      ....
MSV-1 gC 200 TDPHVLWAEAGAGPGAOPPLYSVTG--PLPTORLTIGEVTPATOGMYLWAGR
MSV-2 gF 167 TDPHVLWAEAGAGPGASPRLYSVVGLGRQLIIEELTLETQGMYYWVWGR

      C N NC   C   CC   C   C   N NC
      ....
MSV-1 gC 250 MDSPHEYGTWVRVRFPPSLTLQPHAVMEGQPFKAT--TA--AAYPRNPVE
MSV-2 gF 217 TDRPSAYGTWVRVRFPPSLTIHPHAVLEGQPFKAT--TA--ATYYPGNRAE

      N   N C N C N   C C   C   C   C
      ....
MSV-1 gC 300 FDFEDDROVFNPGQIDTQTHEMPDGF--T--TSTVSTSEAVGGQVPPRTFT
MSV-2 gF 267 FVMFEDGRRVFDPAQIHTQTQENPDGF--T--TSTVSTSAVGGGQVPPRTFT

      C   C   C C C   CCN C
      ....
MSV-1 gC 350 MTWHRDSVTF--SR--N--GLALVLPRTITMEFGYRHYV--TAG--VPEGVTF
MSV-2 gF 317 LTWHRDSVSF--SR--N--GTASVLPRTITMEFTGDHAY--TAG--VPEGVTF

      C   C C CC C NC   C   C   C CCC   N C
      ....
MSV-1 gC 400 WFLGDDPSPAAKSAVTAOES--DHPGLATVRSTLPISYDYSEYI--WLTGYP
MSV-2 gF 367 WFLGDDSSPAEKVAVASOTS--GRPGTATIRSTLPVSYEQEYI--WLAGYP

      N
      ....
MSV-1 gC 450 AGIPVLEMHGSHOPPPRDP--TERQVIEAIEWVGIGIGVLAAGVLVVTAIYV
MSV-2 gF 417 DGIPVLEMHGSHOPPPRDP--TERQVIRAVEGAGIGVAVLVAVVLAGTAVVY

      CCCC NN N NN
      ....
MSV-1 gC 500 VVRTSQSRORHRR
MSV-2 gF 467 LTHASSVRYRRLR

```

B.

```

      C C C   C C CCCC
MSV-2 730bp ORF 1  MAFRASGPATQPLAPRPPPARARYPAVAMIGVGAIVGAFALVALVLP
MSV-1 730bp ORF 1  -----MRARLPAAAWVGVTIIGGVVIAALVLP

      C C C C   C   C CC   C
MSV-2 730bp ORF 50 PRSSWGLCPD--SGWQEFNAGCVAWDPTPVMEHQAVGGCSAPATLIPRAA
MSV-1 730bp ORF 31 SRASWALSPD--SGWHEFNLCISWDPTPVMEHQAVGGCSAPATLIPRAA

      C   CC CC   N   CN   C C N   C
MSV-2 730bp ORF 100 KHLAALTRYOAERSSGYWVWNGDGIRTCLRLVDSVSGIDGFCEEL
MSV-1 730bp ORF 81 KQLAARVQVSARSSGYWVWNGDGIRARLRLVDSVSGIDGFCEELPALRIC

MSV-1 730bp ORF 131 YTPRSPGGFVQFVTSTRNALGLP

```

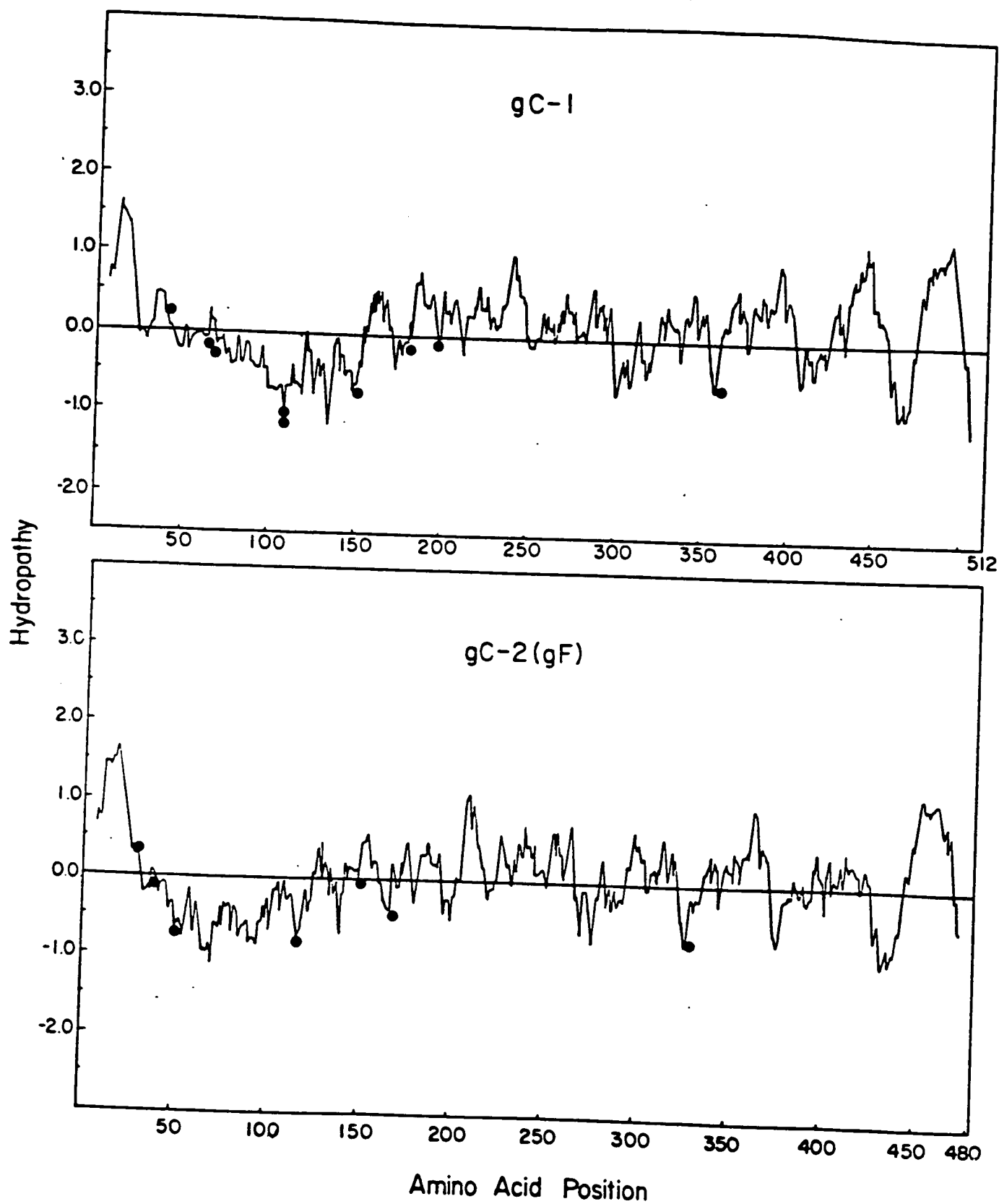


FIGURE 16

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